

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARLRDRGLFVAETKEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKKLKKGVSLLAR....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIIAPQLKNYRWKGINSSGKK	TSGNVLAMTEIEVRERLDAQHIKIKKLKKGKSISFITK....ISH	62
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKL	VNGEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKL	VNGEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1038995	ACCP94 RS05190 WP 057685777.1	MSVARSSIKKQPVDRNVSEIIPFVWEGTDKRGVK	MKGEQQARNANMLRAELRRQGITPTVVKTKPKPLFGA....AGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLTFFHYQGINKAGQK	MEGDIQARSLAIKADLRKQGIVTNKVVKKRKPLFDR....KNK	62
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIAKNSVFTWEGLDLRQGAK	IKGELSGVSPALVKAQLRKQGINPQKVRKKSISLFS....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIG	LKGELPARNMALAKVTLRKQGITTTRTREKRKNIFEG...LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKARQMPTFAYEGVDRKGIVK	IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG...LFFK	64
consensus		* * * * *	! * * * * *	

[illegible][illegible]

